

Product datasheet for SC204970

HARS1 (NM 002109) Human 3' UTR Clone

Product data:

Product Type: 3' UTR Clones

Product Name: HARS1 (NM_002109) Human 3' UTR Clone

Symbol: HARS1

Synonyms: CMT2W; HARS; HRS; USH3B

Mammalian Cell

Selection:

Neomycin

Vector: pMirTarget (PS100062)

ACCN: NM_002109

Insert Size: 382 bp

Insert Sequence: >SC204970 3'UTR clone of NM_002109

The sequence shown below is from the reference sequence of NM_002109. The complete

sequence of this clone may contain minor differences, such as SNPs.

Blue=Stop Codon Red=Cloning site

GGCAAGTTGGACGCCCGCAAGATCCGCGAGATTCTCATTAAGGCCAAGAAGGGCGGAAAGATCGCCGTG

TAACAATTGGCAGAGCTCAGAATTCAAGCGATCGCC

AATTGCCAAAGGTCCAATAAAATGCCTCAACCACGGA

ACGCGTAAGCGGCCGCGCATCTAGATTCGAAGAAAATGACCGACCAAGCGACGCCCAACCTGCCATCA

CGAGATTTCGATTCCACCGCCGCCTTCTATGAAAGG

Restriction Sites: Sgfl-Mlul

OTI Disclaimer: Our molecular clone sequence data has been matched to the sequence identifier above as a

point of reference. Note that the complete sequence of this clone is largely the same as the

reference sequence but may contain minor differences, e.g., single nucleotide

polymorphisms (SNPs).

Components: The cDNA clone is shipped in a 2-D bar-coded Matrix tube as 10 ug dried plasmid DNA. The

package also includes 100 pmols of both the corresponding 5' and 3' vector primers in

separate vials.



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RefSeq: <u>NM 002109.6</u>

Summary: Aminoacyl-tRNA synthetases are a class of enzymes that charge tRNAs with their cognate

amino acids. The protein encoded by this gene is a cytoplasmic enzyme which belongs to the class II family of aminoacyl-tRNA synthetases. The enzyme is responsible for the synthesis of histidyl-transfer RNA, which is essential for the incorporation of histidine into proteins. The gene is located in a head-to-head orientation with HARSL on chromosome five, where the homologous genes share a bidirectional promoter. The gene product is a frequent target of autoantibodies in the human autoimmune disease polymyositis/dermatomyositis. Several transcript variants encoding different isoforms have been found for this gene. [provided by

RefSeq, Apr 2012]

Locus ID: 3035 MW: 14.4